

Supplementary Table 1 Percent sequence identity within 24 conserved miRNA families under salt stress

Sl. no.	miRNA family	No. miRNAs	Min (% identity)	Max (% identity)	Average (% identity)
1	miR156	11	88.88	100	95.03
2	miR160	13	90.47	100	98.57
3	miR164	11	85.71	100	95.76
4	miR166	11	85.00	100	96.01
5	miR167	11	90.00	100	95.97
6	miR169	19	71.42	100	89.19
7	miR171	12	84.21	100	94.09
8	miR172	12	95.00	100	98.01
9	miR319	9	89.47	100	96.48
10	miR393	9	90.47	100	97.75
11	miR394	7	95.00	100	97.85
12	miR395	15	90.47	100	96.91
13	miR396	6	72.22	100	94.32
14	miR390	7	84.21	100	94.81
15	miR399	4	85.71	100	92.85
16	miR168	5	90.47	100	95.67
17	miR482	3	85.00	100	87.12
18	miR159	3	95.00	100	96.74
19	miR530	2	100	100	100
20	miR398	2	94.44	94.44	94.44
21	miR408	2	100	100	100
22	miR528	2	100	100	100
23	miR1551	2	85.00	85.00	85.00
24	miR2111	2	100	100	100

